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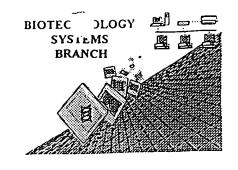
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## RAW SEQUENCE LISTING ERROR REPORT



0590

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/892,864Source: 01PEDate Processed by STIC: 7/12/2000

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
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- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 c-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 c-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

#### Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

DATE: 07/12/2001

TIME: 09:44:33

#### OIPE

Input Set : A:\209524US0CONT.txt Output Set: N:\CRF3\07122001\I892864.raw 3 <110> APPLICANT: YOKOYAMA, Keiichi ONO, Kunio EJIMA, Daisuke 7 <120> TITLE OF INVENTION: PROCESS FOR PRODUCING TRANSGLUTAMINASE 9 <130> FILE REFERENCE: 209524US0CONT C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/892,864 C--> 11 <141> CURRENT FILING DATE: 2001-06-28 **Does Not Comply** 11 <150> PRIOR APPLICATION NUMBER: PCT/JP99/07250 Corrected Diskette Needed 12 <151> PRIOR FILING DATE: 1999-12-24 14 <150> PRIOR APPLICATION NUMBER: JP 10-373131 15 <151> PRIOR FILING DATE: 1998-12-28 17 <160> NUMBER OF SEQ ID NOS: 58 19 <170> SOFTWARE: PatentIn version 3.1 21 <210> SEQ ID NO: 1 22 <211> LENGTH: 1519 23 <212> TYPE: DNA 24 <213> ORGANISM: Artificial Sequence 26 <220> FEATURE: 27 <223> OTHER INFORMATION: Synthetic DNA 29 <220> FEATURE: 30 <221> NAME/KEY: CDS 31 <222> LOCATION: (87)..(1082) 32 <223> OTHER INFORMATION: 35 <400> SEQUENCE: 1 36 ttcccctgtt gacaattaat catcgaacta gttaactagt acgcaagttc acgtaaaaag 38 ggtatcgatt agtaaggagg tttaaa atg gat tct gac gat cgt gtt act cca 113 39 Met Asp Ser Asp Asp Arg Val Thr Pro 40 42 cca gct gaa cca ctg gat cgt atg cca gat cca tat cgt cca tct tat 161 43 Pro Ala Glu Pro Leu Asp Arg Met Pro Asp Pro Tyr Arg Pro Ser Tyr 15 20 209 46 ggt cgt gct gaa act gtt gtt aat aat tat att cgt aaa tgg caa caa 47 Gly Arg Ala Glu Thr Val Val Asn Asn Tyr Ile Arg Lys Trp Gln Gln 30 35 257 50 gtt tat tct cat cgt gat ggt cgt aaa caa caa atg act gaa gaa caa 51 Val Tyr Ser His Arg Asp Gly Arg Lys Gln Gln Met Thr Glu Glu Gln 50 52 305 54 cgt gaa tgg ctg tct tat ggt tgc gtt ggt gtt act tgg gtt aac tct 55 Arg Glu Trp Leu Ser Tyr Gly Cys Val Gly Val Thr Trp Val Asn Ser 58 ggt cag tat ccg act aac cgt ctg gca ttc gct tcc ttc gat gaa gat 353 59 Gly Gln Tyr Pro Thr Asn Arg Leu Ala Phe Ala Ser Phe Asp Glu Asp 62 cgt ttc aag aac gaa ctg aag aac ggt cgt ccg cgt tct ggt gaa act 401 63 Arg Phe Lys Asn Glu Leu Lys Asn Gly Arg Pro Arg Ser Gly Glu Thr 64 90 100 66 cgt gct gaa ttc gaa ggt cgt gtt gct aag gaa tcc ttc gat gaa gag 449

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/892,864



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Output Set: N:\CRF3\07122001\1892864.raw

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71 Lys Gly Phe Gln Arg Ala Arg Glu Val Ala Ser Val Met Asn Arg Ala	
72 125 130 135 74 sta gag and gat gat gat gat tot gat tog sta gat and gat and gat	EAE
74 cta gag aac gct cat gat gaa tct gct tac ctg gat aac ctg aag aag 75 Leu Glu Asn Ala His Asp Glu Ser Ala Tyr Leu Asp Asn Leu Lys Lys	545
76 140 145 150	
78 gaa ctg gct aac ggt aac gat gct ctg cgt aac gaa gat gct cgt tct	593
79 Glu Leu Ala Asn Gly Asn Asp Ala Leu Arg Asn Glu Asp Ala Arg Ser	333
80 155 160 165	
82 ccg ttc tac tct gct ctg cgt aac act ccg tcc ttc aaa gaa cgt aac	641
83 Pro Phe Tyr Ser Ala Leu Arg Asn Thr Pro Ser Phe Lys Glu Arg Asn	
84 170 175 180 185	
86 ggt ggt aac cat gat ccg tct cgt atg aaa gct gtt atc tac tct aaa	689
87 Gly Gly Asn His Asp Pro Ser Arg Met Lys Ala Val Ile Tyr Ser Lys	
88 190 195 200	727
90 cat ttc tgg tct ggt cag gat aga tct tct tct gct gat aaa cgt aaa 91 His Phe Trp Ser Gly Gln Asp Arg Ser Ser Ser Ala Asp Lys Arg Lys	737
92 205 210 215	
94 tac ggt gat ccg gat gca ttc cgt ccg gct ccg ggt act ggt ctg gta	785
95 Tyr Gly Asp Pro Asp Ala Phe Arg Pro Ala Pro Gly Thr Gly Leu Val	, 00
96 220 225 230	
98 gac atg tct cgt gat cgt aac atc ccg cgt tct ccg act tct ccg ggt	833
99 Asp Met Ser Arg Asp Arg Asn Ile Pro Arg Ser Pro Thr Ser Pro Gly	
100 235 240 245	
102 gaa ggc ttc gtt aac ttc gat tac ggt tgg ttc ggt gct cag act gaa	881
103 Glu Gly Phe Val Asn Phe Asp Tyr Gly Trp Phe Gly Ala Gln Thr Glu	
104 250 . 255 . 260 . 265	000
106 gct gat gct gat aag act gta tgg acc cat ggt aac cat tac cat gct 107 Ala Asp Ala Asp Lys Thr Val Trp Thr His Gly Asn His Tyr His Ala	929
108 270 275 280	
110 ccg aac ggt tct ctg ggt gct atg cat gta tac gaa tct aaa ttc cgt	977
111 Pro Asn Gly Ser Leu Gly Ala Met His Val Tyr Glu Ser Lys Phe Arg	3.,
112 285 290 295	
114 aac tgg tct gaa ggt tac tct gac ttc gat cgt ggt gct tac gtt atc	1025
115 Asn Trp Ser Glu Gly Tyr Ser Asp Phe Asp Arg Gly Ala Tyr Val Ile	
116 300 305 310	
118 acc ttc att ccg aaa tct tgg aac act gct ccg gac aaa gtt aaa cag	1073
119 Thr Phe Ile Pro Lys Ser Trp Asn Thr Ala Pro Asp Lys Val Lys Gln	
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124 330	
126 actaaaatag acatatetta tattatgtga ttttgtgaca tttcctagat gtgaggtgga	1182
128 ggtgatgtat aaggtagatg atgateetet aegeeggaeg categtggee ggeateaeeg	1242
130 gcgccacagg tgcggttgct ggcgcctata tcgccgacat caccgatggg gaagatcggg	1302
132 ctcgccactt cgggctcatg agcgcttgtt tcggcgtggg tatggtggca ggccccgtgg	1362
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RAW SEQUENCE LISTING DATE: 07/12/2001 PATENT APPLICATION: US/09/892,864 TIME: 09:44:33

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		2> T			<i>.</i> .											i		DAIA
					Art	ific	י ובו	Seans	ance			1	41	( )	ς Λ	10+	a	DON
	4 <213> ORGANISM: Artificial Sequence +his is 7407 456 <220> FEATURE:																	
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						Arg	Val	Thr	Dro	Dro								
152		дор	Ser	тэр	дэр 5	Arg	Val	1111	rio	10	пта	GIU	FIO	neu	15	Arg		
		Dro	Δen	Dro	-	Arg	Pro	Sar	Tur		Ara	ЛΊэ	Glu	Thr		Wal		
156	1100	110	изъ	20	ı yı	Arg	110	561	25	Оту	Arg	AIQ	GIU	30	val	Vai		
	Aen	Aen	ጥኒፖ		Δνα	Lys	Ψrn	Gln		V = 1	Тиг	Sar	Hic		λευ	Clv	•	•
160	ASII	N311	35	116	Arg	БУЗ	тър	40	GIII	Val	ıyı	PET	45	Arg	тэр	Gry		
	Δra	Luc		Gln	Mot	Thr	Glu		Gln	Δrα	Glu	Trn		Sar	Тиг	Gly		
164	Arg	50	OIII	GLII	1100	1111	55	Giu	GIII	AL Y	Giu	60	пец	Ser	ıyı	СТУ		
	Cue		Glv	V = 1	Thr	Trp		Aen	Sar	Gly	Gln		Dro	Thr	Λen	Λrα		
168		val	оту	Val	1111	70	Val	. non	Ser	СТУ	75	тут	110	1111	ASII	80		
		Δla	Phe	Δla	Sar	Phe	Aen	Glu	Aen	Δrα		Luc	Δe'n	Glu	Len			
172	пси	711 U	1110	1114	85	LIIC	тор	OIU	тор	90	LIIC	БУЗ	non.	Olu	95	цуз		
	Aen	Glv	Ara	Pro		Sor	Glv	Glu	Thr		Δla	Glu	Pho	Glu		Arg		•
176	ASII	OLY	rig	100	Arg	SCI	СТУ	Giu	105	nrg	лта	Olu	1116	110	СТУ	Arg.	•	
	Val	Δla	Lus		Ser	Phe	Asn	Glu		T.ve	Glv	Phe	Gln		Δla	Δra		
180	VUI	mu	115	Olu	JUL	1110	пор	120	Olu	цуз	OLY	1110	125	my	mu	my		
	Glu	Val		Ser	Va 1	Met	Δsn		Δla	T.e.1	Glu	Asn	-	His	Δsn	Glu		
184	OIU	130	1114	DCI	VUI	1100	135	711.9	711 U	цец	OIU	140	71±Q	1115	тор	O.L.u		•
	Ser		Ψur	Len	Asn	Asn		T.vs	T.vs	Glu	T.e.i		Δsn	Glv	Asn	Asn		
	145	1114	- y -	ьçц	nop	150	цси	Lyo	БуЗ	O L u	155	1114	11011	O <sub>T</sub> y	11011	160		
		T.em	Ara	Asn	Glu	Asp	Ala	Ara	Ser	Pro		Tur	Ser	Δla	T.e.ii			
192	1114	ncu.	1119	11011	165	1150	1114	1119	UC L	170	1110	+ y +	DCI	1114	175	111.9	•	
	Asn	Thr	Pro	Ser		Lys	Glu	Ara	Asn		Glv	Asn	His	Asp	•	Ser		
196				180		2,0		•••	185	011	01,			190		501		
	Ara	Met	Lvs		Val	Ile	Tvr	Ser		His	Phe	Trp	Ser		Gln	Asp		
200	5		195				- 1 -	200	-1-				205	1		1101		
	Ara	Ser		Ser	Ala	Asp	Lvs		Lvs	Tvr	Glv	Asp		Asp	Ala	Phe		
204	_	210				•	215	,	-	-	-	220		•				
207	Arq		Ala	Pro	Gly	Thr	Gly	Leu	Val	Asp	Met	Ser	Arg	Asp	Arq	Asn		
	225				-	230	-			•	235		,	•	,	240		
		Pro	Arg	Ser	Pro	Thr	Ser	Pro	Gly	Glu		Phe	Val	Asn	Phe			
212					245				-	250	-				255	-		
	Tyr	Gly	Trp	Phe	Gly	Ala	Gln	Thr	Glu	Ala	Asp	Ala	Asp	Lys	Thr	Val		
216	-	-	-	260	-				265		_		_	270				
	Trp	Thr	His	Gly	Asn	His	Tyr	His	Ala	Pro	Asn	Gly	Ser	Leu	Gly	Ala		
220	-		275	-			-	280				-	285		-			
223	Met	His	Val	Tyr	Glu	Ser	Lys	Phe	Arg	Asn	Trp	Ser	Glu	Gly	Tyr	Ser		
224		290		-			295		_		-	300		-	-			
227	Asp	Phe	Asp	Arg	Gly	Ala	Tyr	Val	Ile	Thr	Phe	Ile	Pro	Lys	Ser	Trp		
228	305					310					315					320		



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Output Set: N:\CRF3\07122001\1892864.raw

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VERIFICATION SUMMARY PATENT APPLICATION: US/09/892,864

DATE: 07/12/2001 TIME: 09:44:34

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L:11 M:270 C: Current Application Number differs, Replaced Current Application No L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date